

FIG. 1

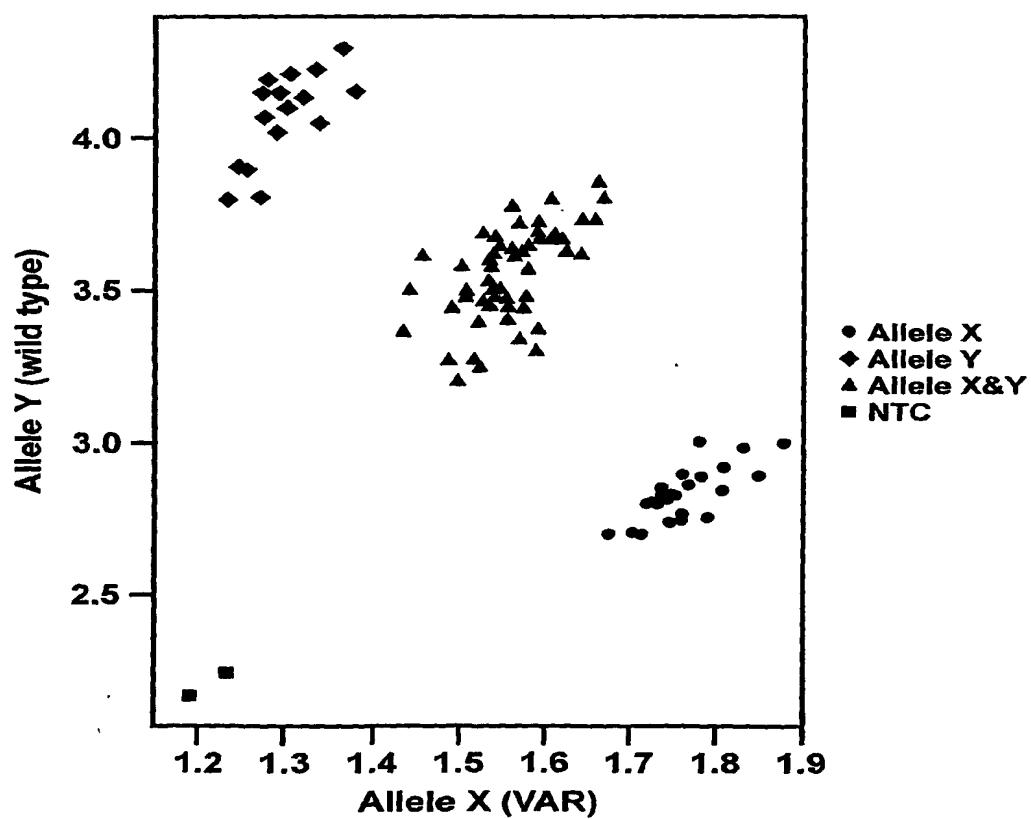
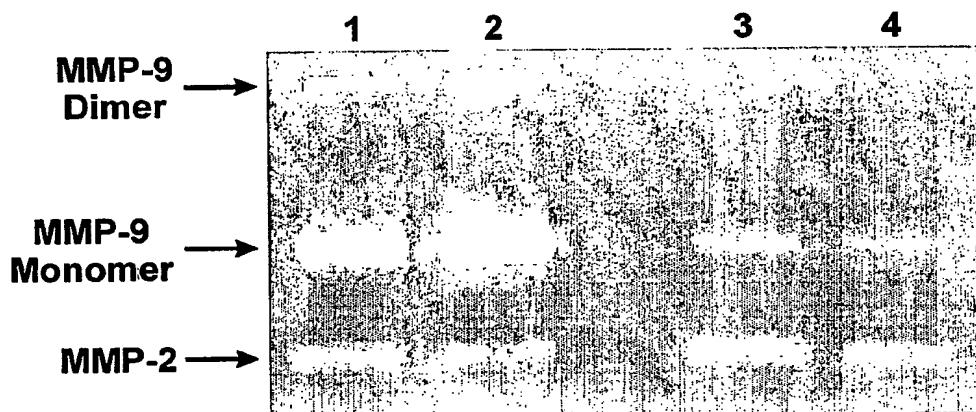
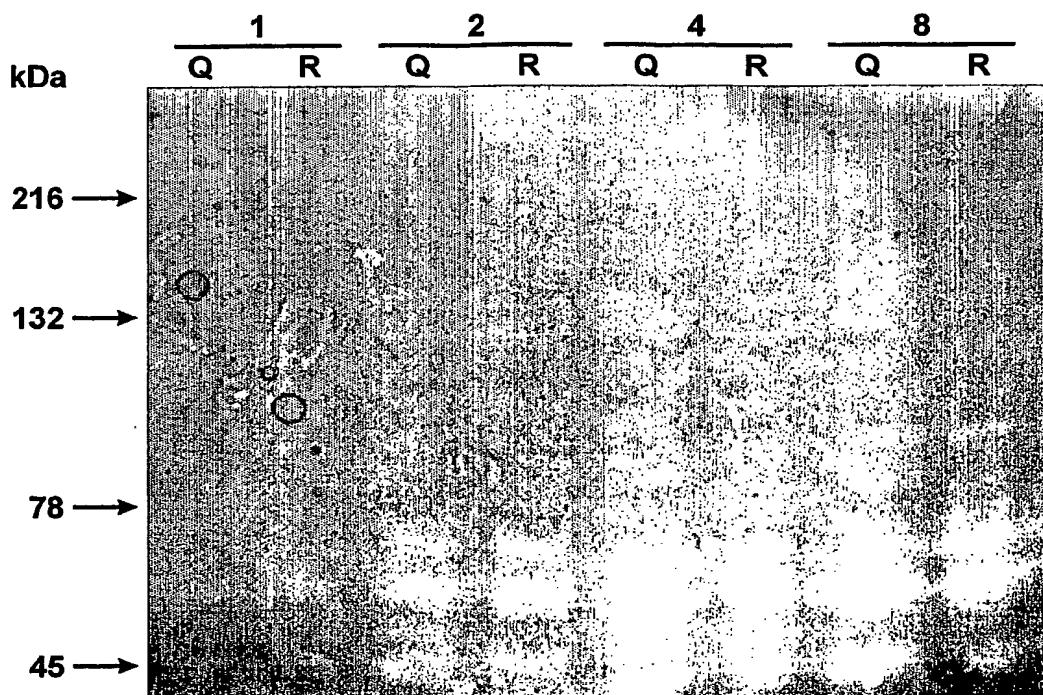


FIG. 2

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Polymorphism	Primers	Probes
C/T-1562	5'-GCCTGGCACATAGTAGGCC (S) (SEQ ID NO:14)	No SphI site (C)
	5'-CTTCCTAGCCAGCCGGCATC (AS) (SEQ ID NO:15)	SphI site 300+260 bp (T)
A/G Gln279Arg	5'-TCTCCCCCTTCCCACATC (S) (SEQ ID NO:3)	5'-CTCTACACCC <u>GGG</u> ACGGCAATG (SEQ ID NO:1)
	5'-TGTGCTGTCTCCGCCCTCT (AS) (SEQ ID NO:4)	5'ACTCTACACCC <u>AGG</u> ACGGCAATGC (SEQ ID NO:2)
CA repeat	5'-6FAM- AAGCCCTTCTCATGCTGGTGC (SEQ ID NO:12)	Fragment Analysis
	5'-TGACAGGCAAGTGCTGACTCAG (SEQ ID NO:13)	

**FIG. 3****FIG. 4****BEST AVAILABLE COPY**



**FIG. 5**

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